

Sequence alignment of protein 40570013.q across various species. The alignment shows sequence conservation and divergence across three main regions: 1000-990, 1200-1190, and 1210-1250 amino acid positions. The alignment includes color-coded regions for hydrophobicity and secondary structure predictions.

1000

40570013.q, 40570012.q, 73710002.q, 73710004.q, AK064052.q, 15240720.p, 43220007.q, AK060260.q, 15225286.p, 2phk_PhK, lias_TGFbR, lias_IRK

1200

40570013.q, 40570012.q, 73710002.q, 73710004.q, AK064052.q, 15240720.p, 43220007.q, AK060260.q, 15225286.p, 2phk_PhK, lias_TGFbR, lias_IRK

1210

40570013.q, 40570012.q, 73710002.q, 73710004.q, AK064052.q, 15240720.p, 43220007.q, AK060260.q, 15225286.p, 2phk_PhK, lias_TGFbR, lias_IRK